

# 20.109

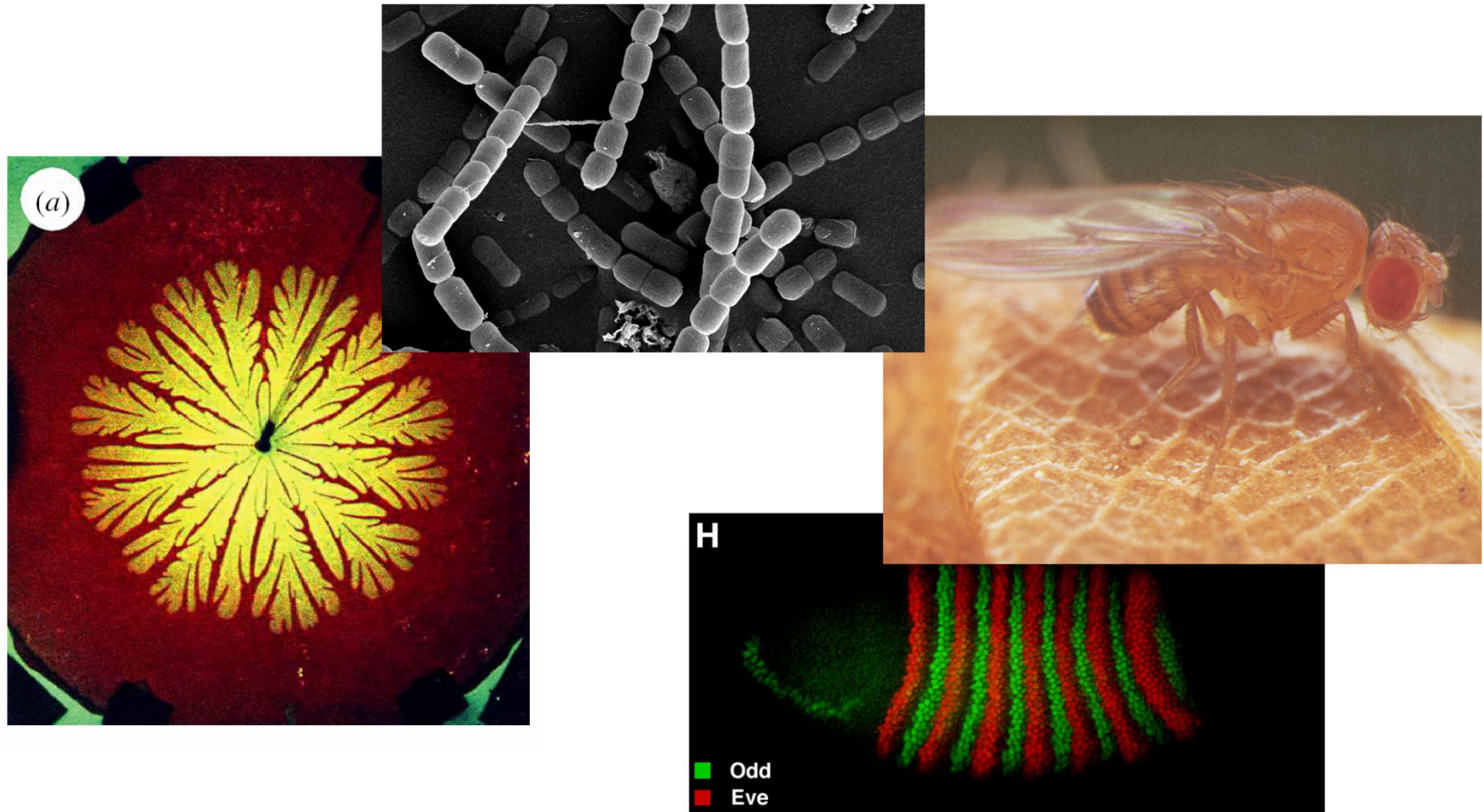
## Synthetic Biology Module

### Lecture #8

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Department of Biological Engineering  
MIT

# Natural Pattern Formation



- Robust global behavior from unreliable parts
- Repeated network motifs. Same molecules used by different species, different stages of development.

# Programmed pattern formation in bacteria

High level behavior:

“Ring-like spatial patterns”

Bio-program:

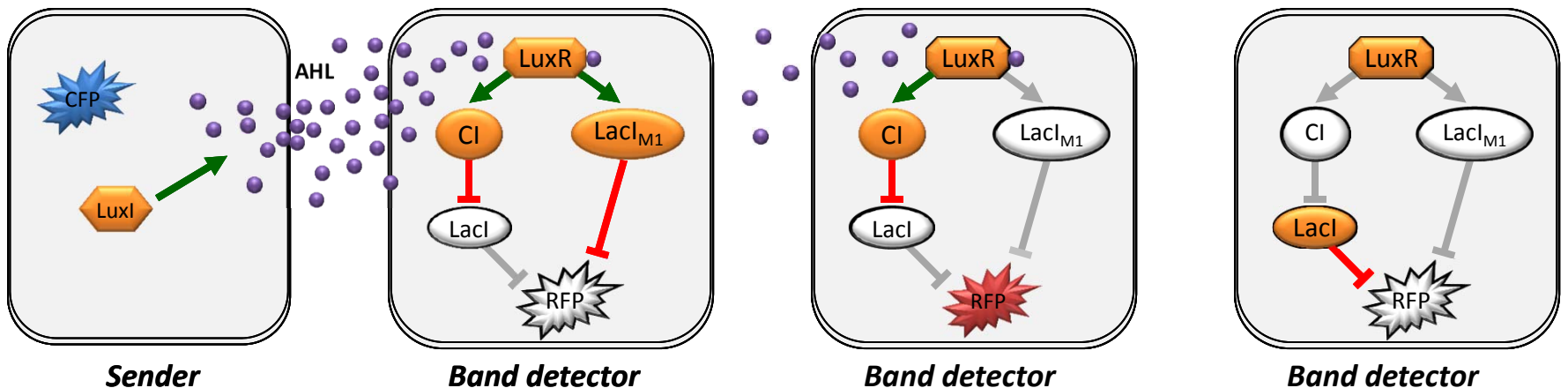
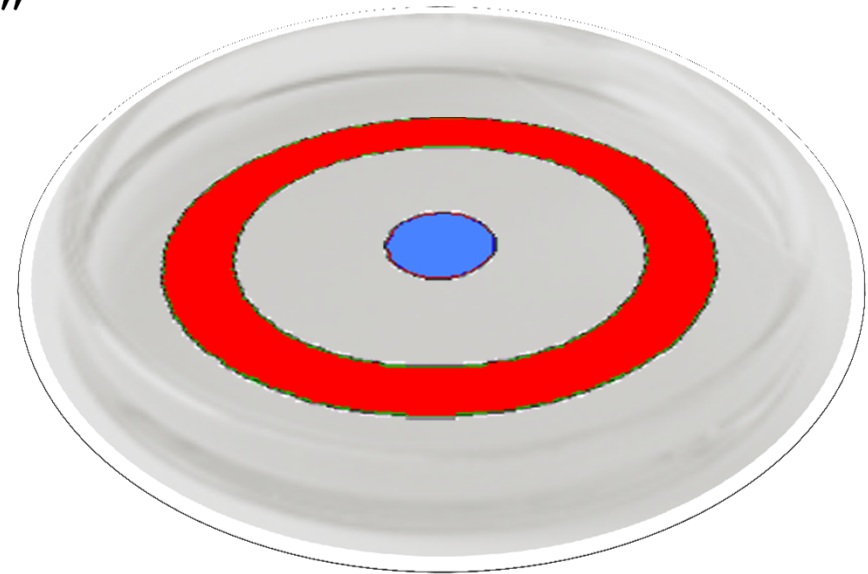
- Cell type: *sender*  
send (AHL)
- Cell type: *receiver*  
input: AHL

case (AHL)

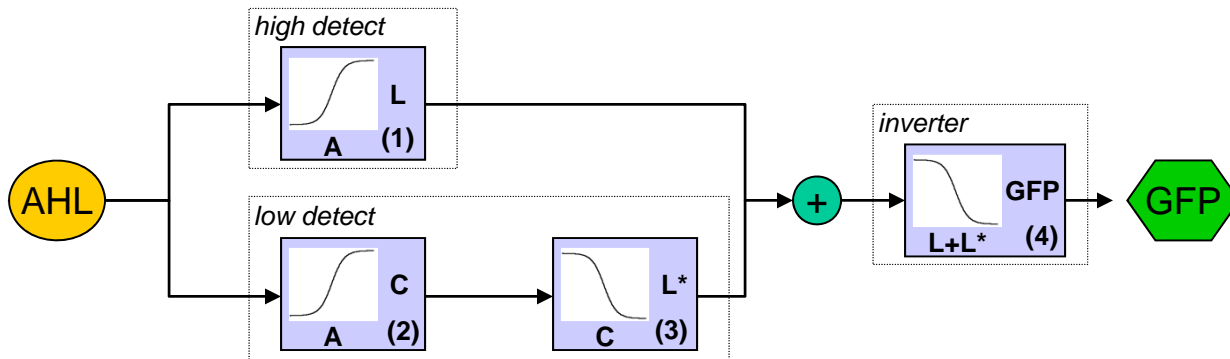
low: RFP = off

med: RFP = on

high: RFP = off

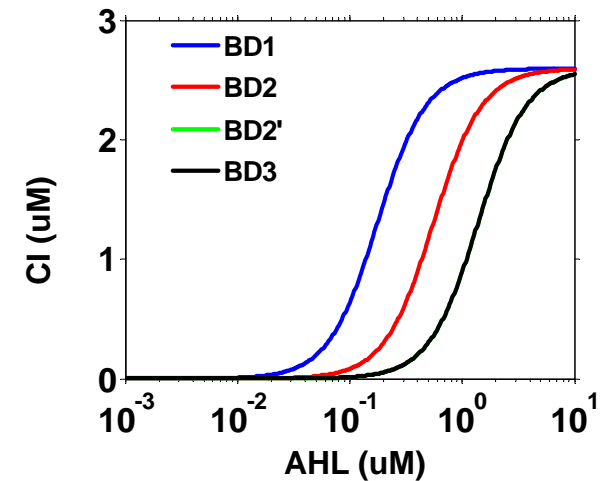
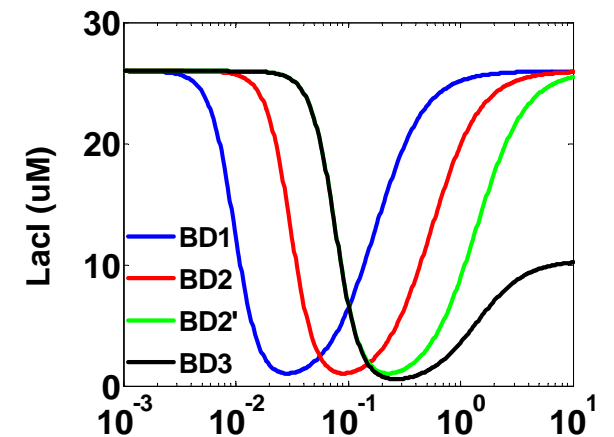
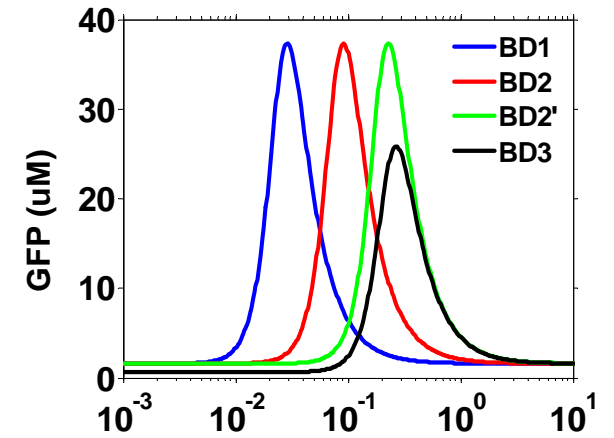


# Band Detect Model

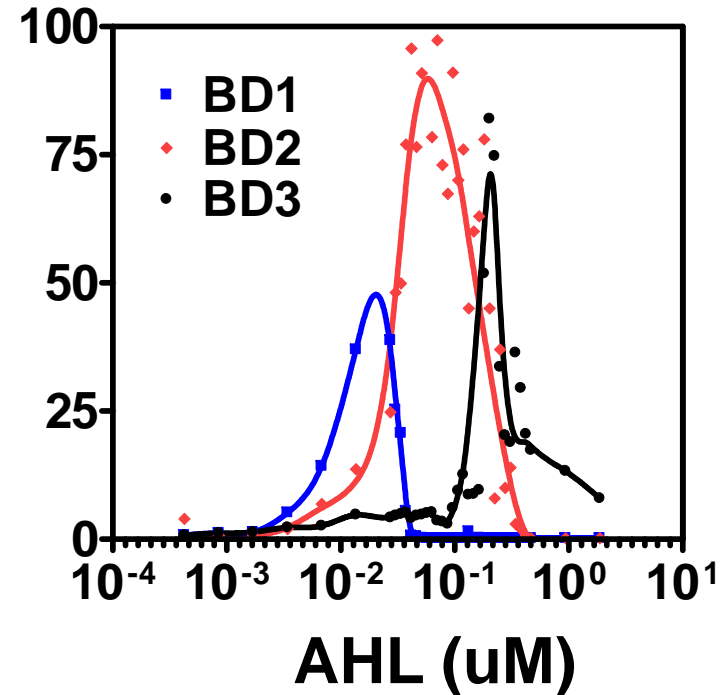
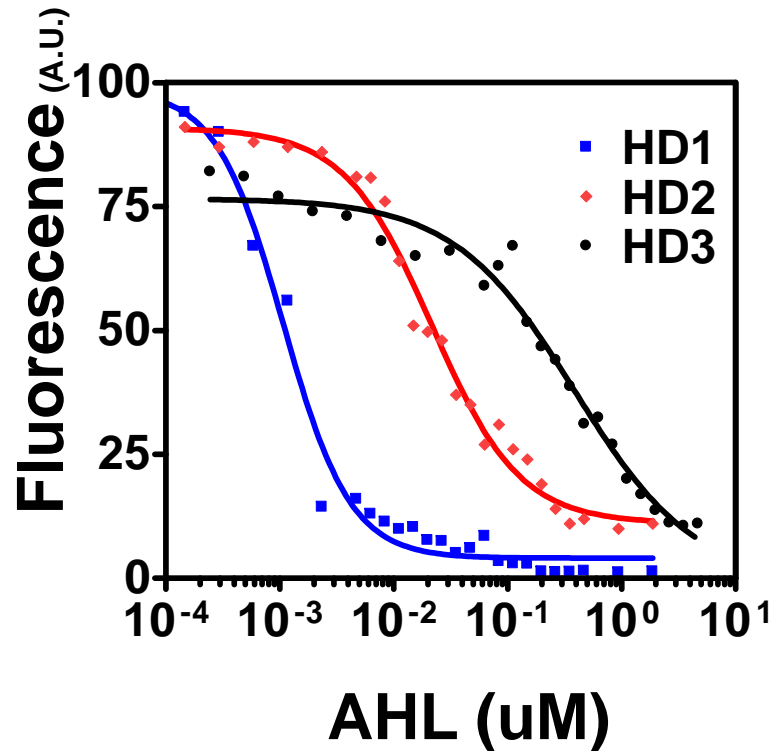


$$\begin{aligned}\frac{dL}{dt} &= \frac{\alpha_L \cdot A^{\eta_A}}{A^{\eta_A} + \beta_A^{\eta_A}} - \gamma_L \cdot L \\ \frac{dC}{dt} &= \frac{\alpha_C \cdot A^{\eta_A}}{A^{\eta_A} + \beta_A^{\eta_A}} - \gamma_C \cdot C \\ \frac{dL^*}{dt} &= \frac{\alpha_{L^*} \cdot \beta_C^{\eta_C}}{\beta_C^{\eta_C} + C^{\eta_C}} - \gamma_{L^*} \cdot L^* \\ \frac{dG}{dt} &= \frac{\alpha_G \cdot \beta_L^{\eta_L}}{\beta_L^{\eta_L} + (L + L^*)^{\eta_L}} - \gamma_G \cdot G\end{aligned}$$

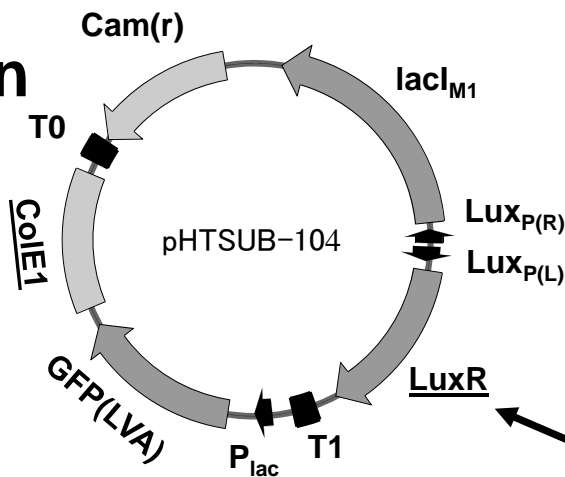
- BD1 – Hypersensitive LuxR
- BD2 – Wildtype LuxR
- BD2' – Reduced plasmid copy number for LuxR
- BD3 – Reduced plasmid copy number for LuxR, LacI<sub>M1</sub>, GFP



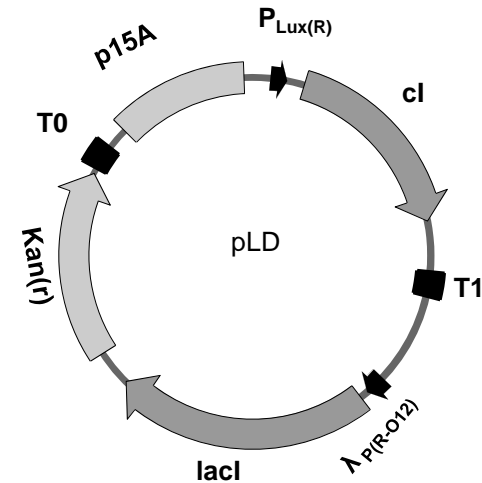
# Experimental Dosage Response



**HD3  
Mutation**

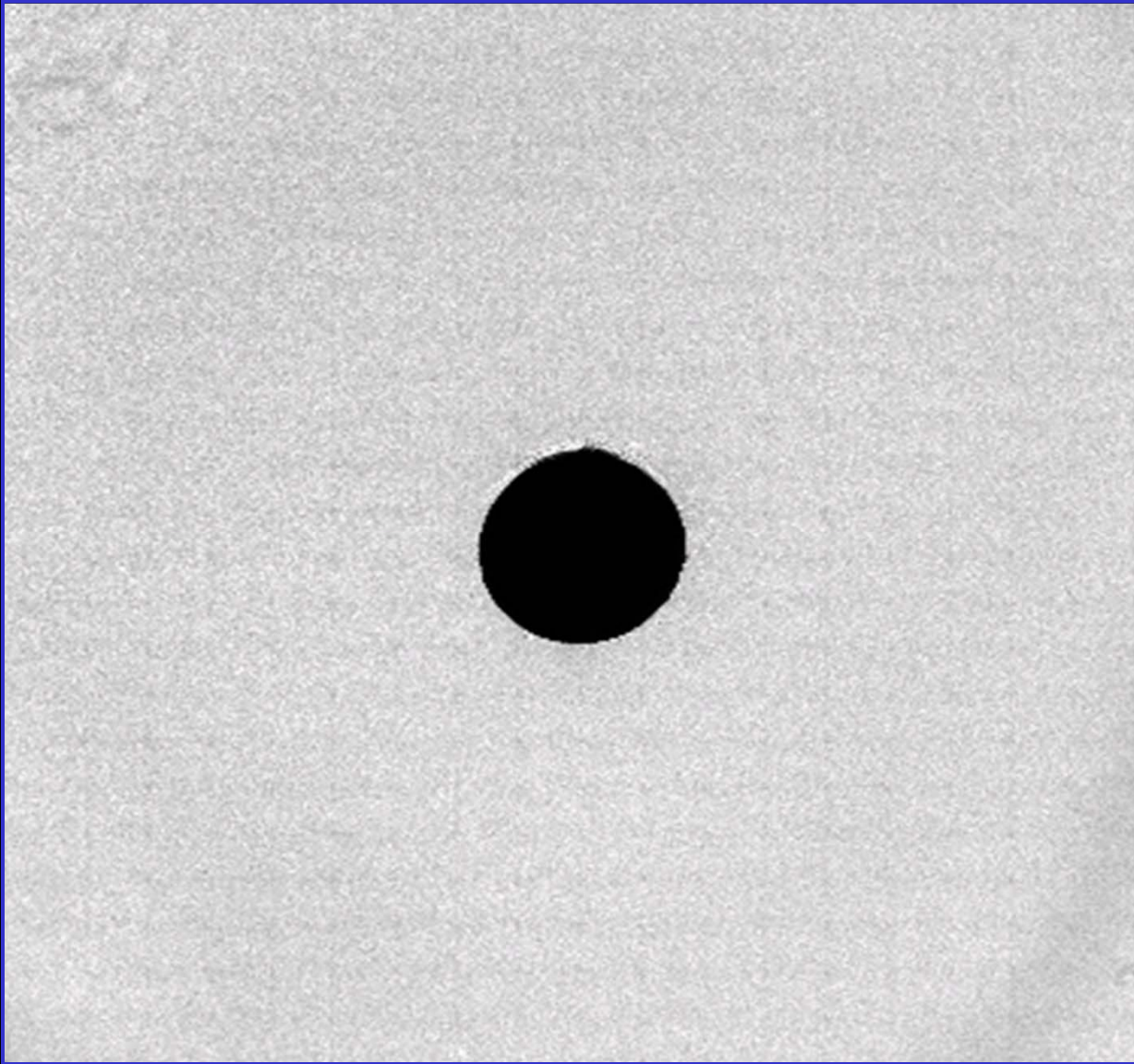


**HD1  
Mutation**

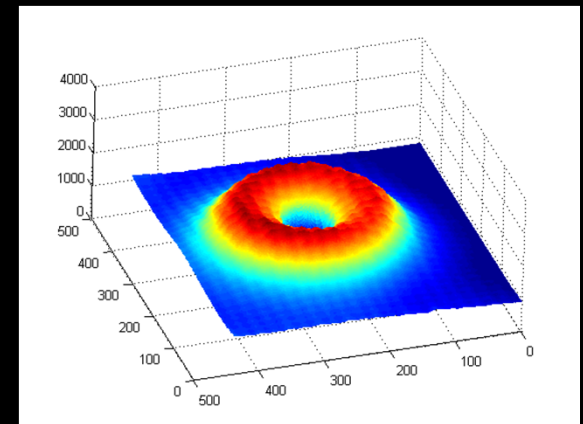
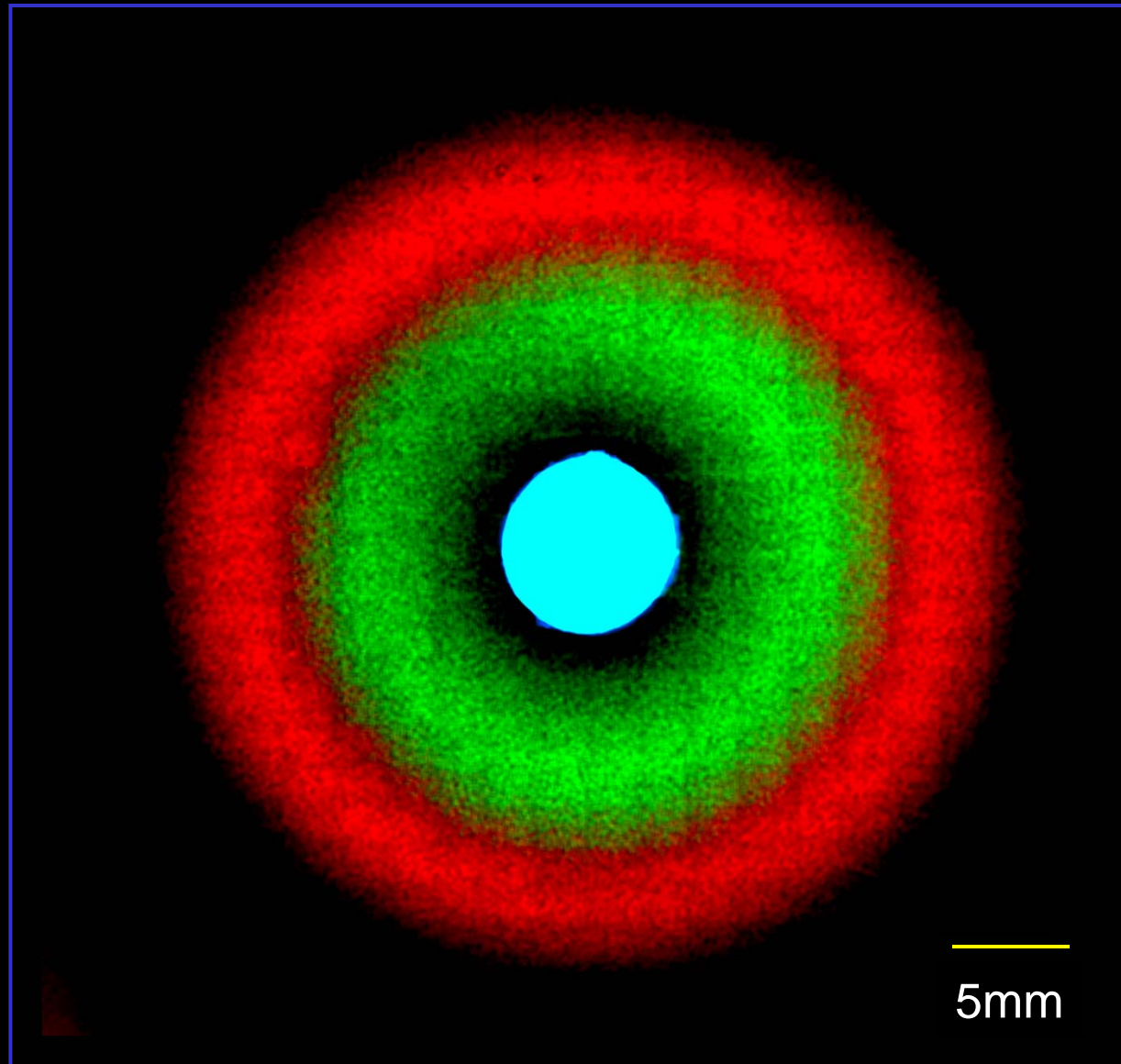




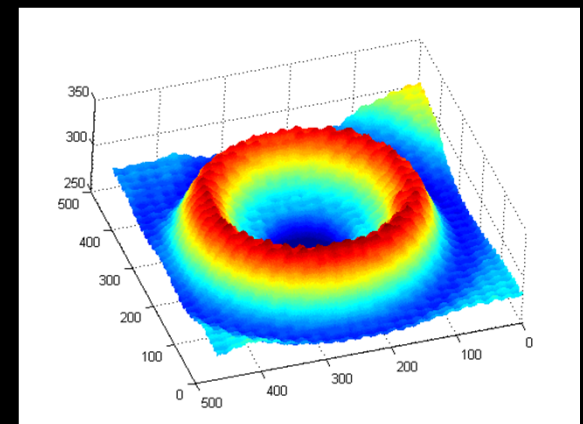
# Bullseye with BD2-Red / BD3-GFP



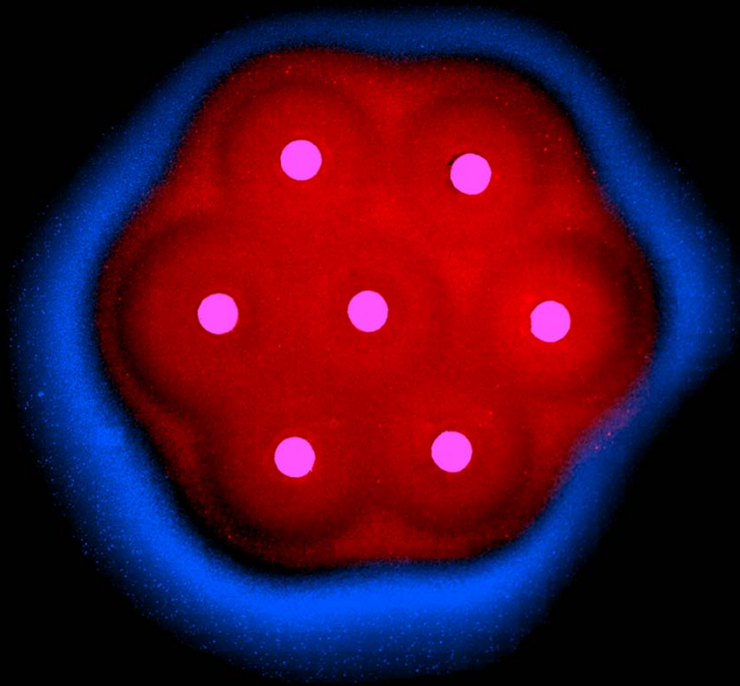
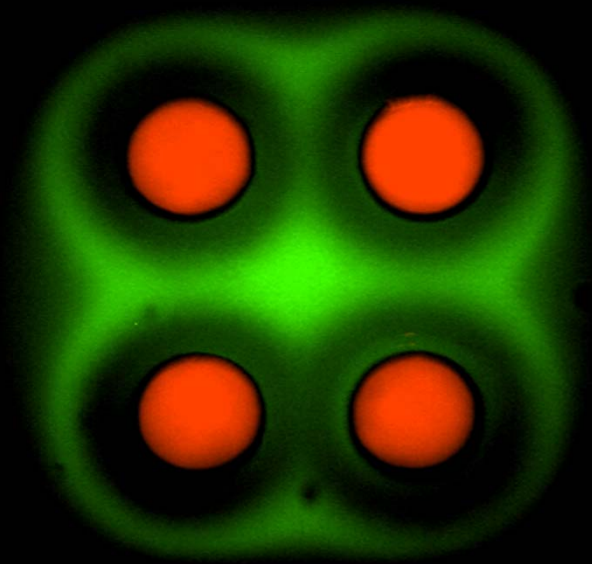
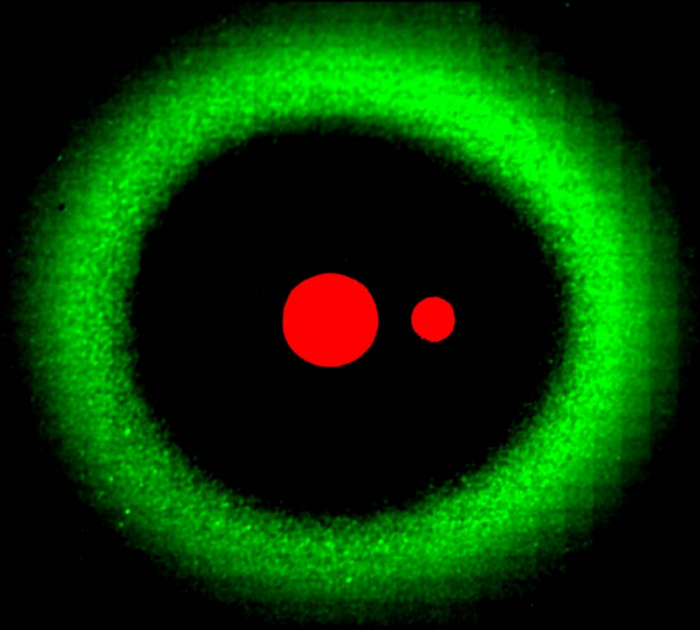
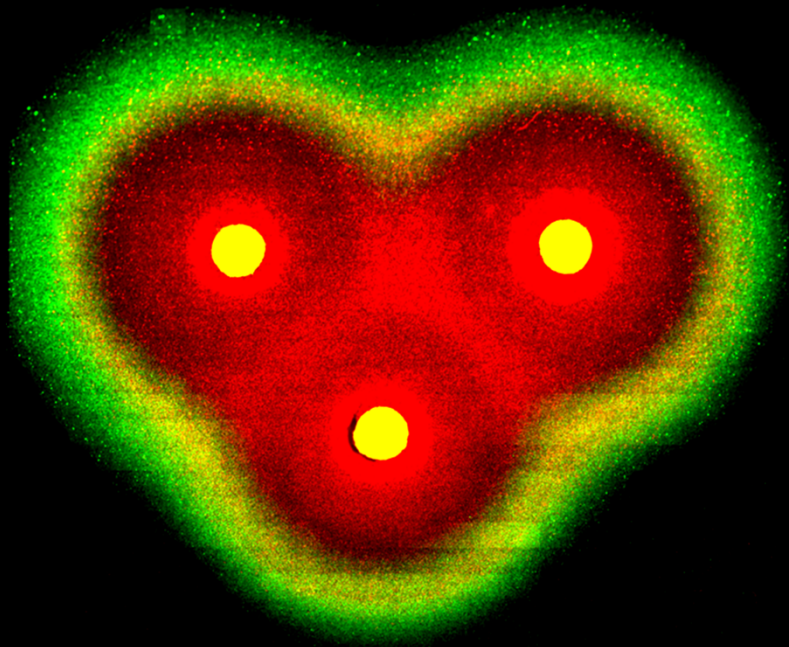
# Bullseye with BD2-Red / BD3-GFP



Green



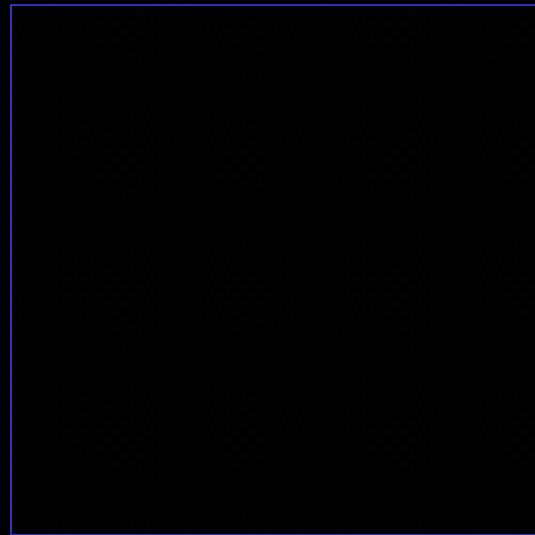
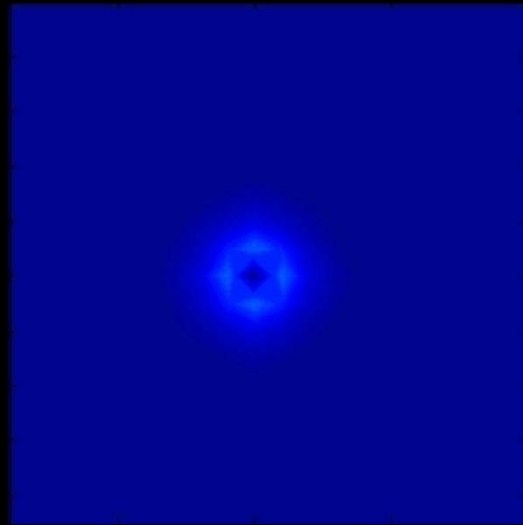
Red



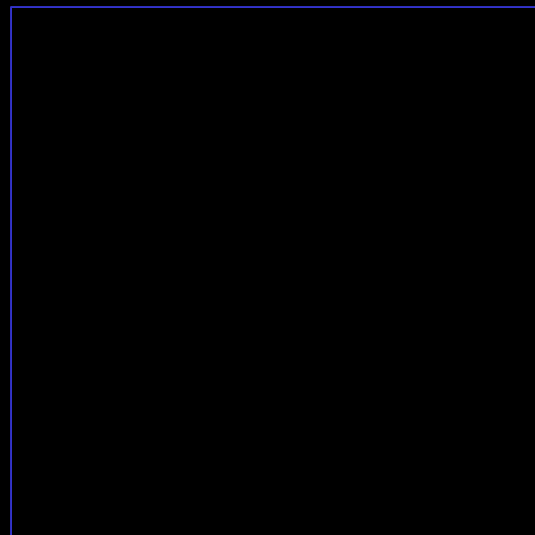
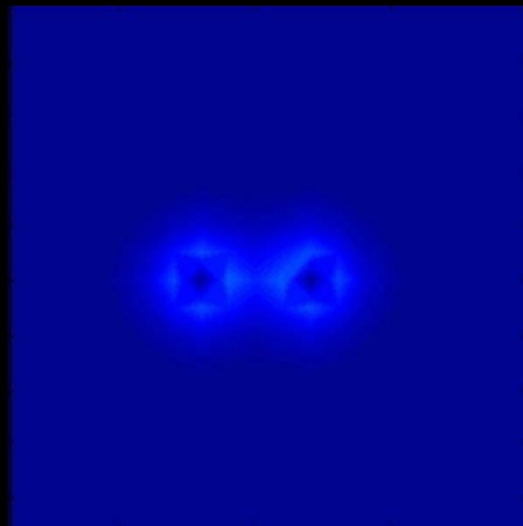


# Spatiotemporal Simulation

*One sender*



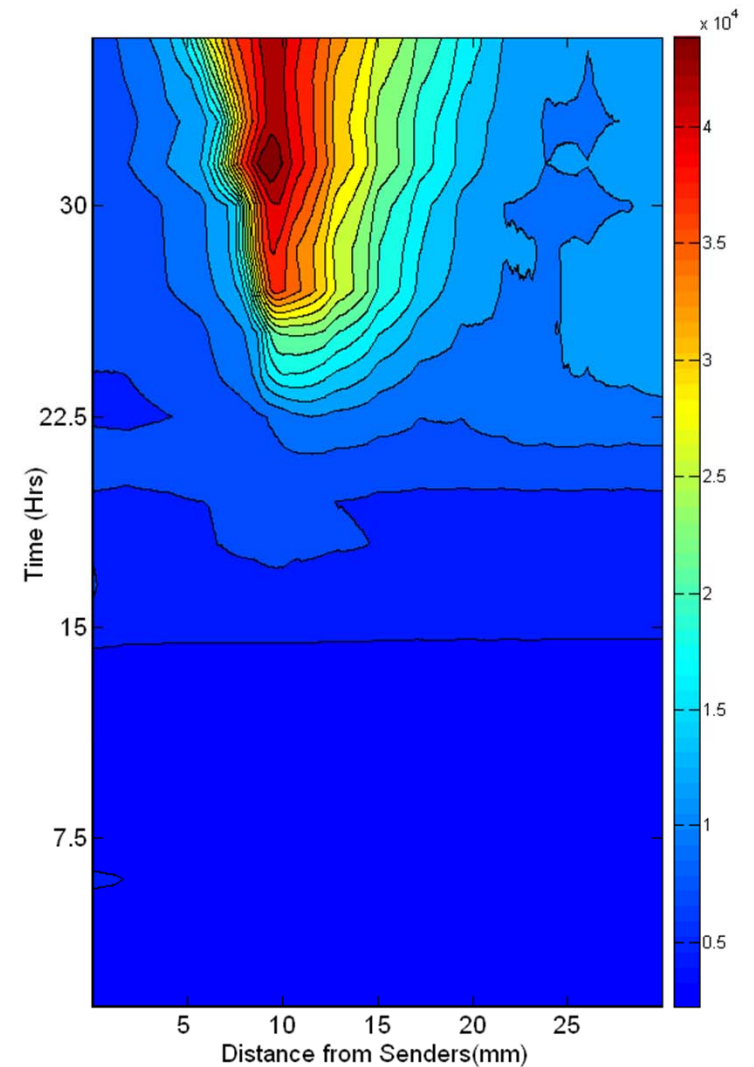
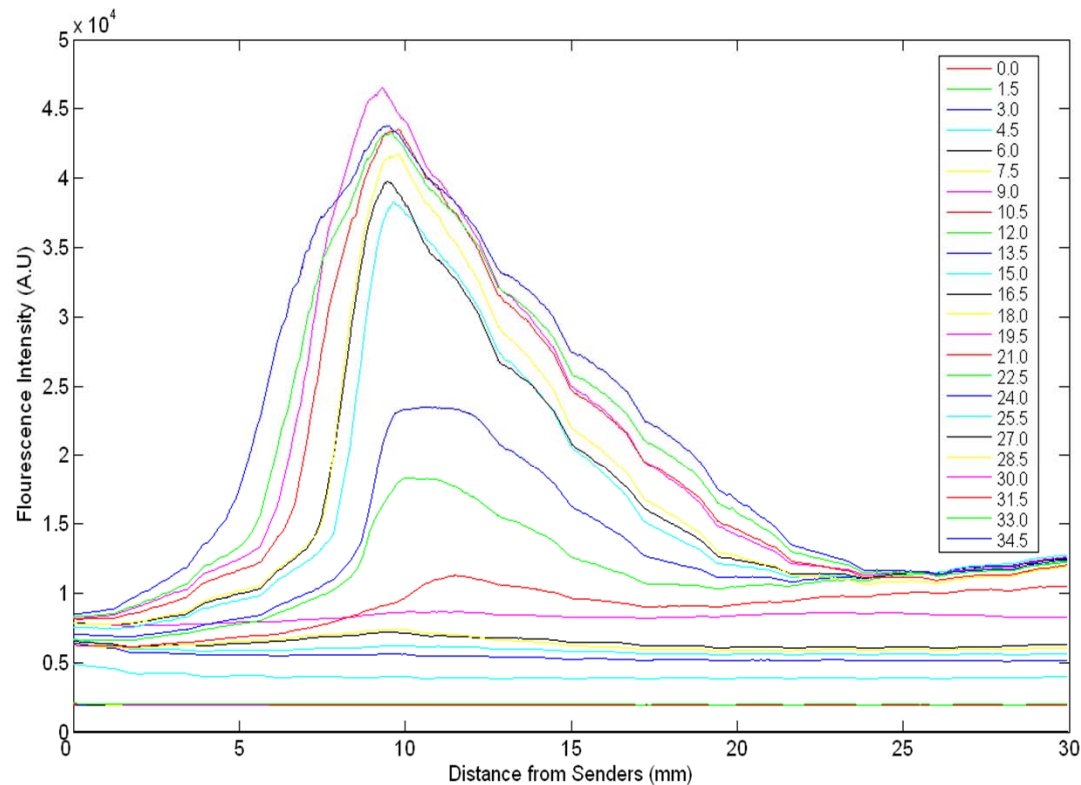
*Two senders*



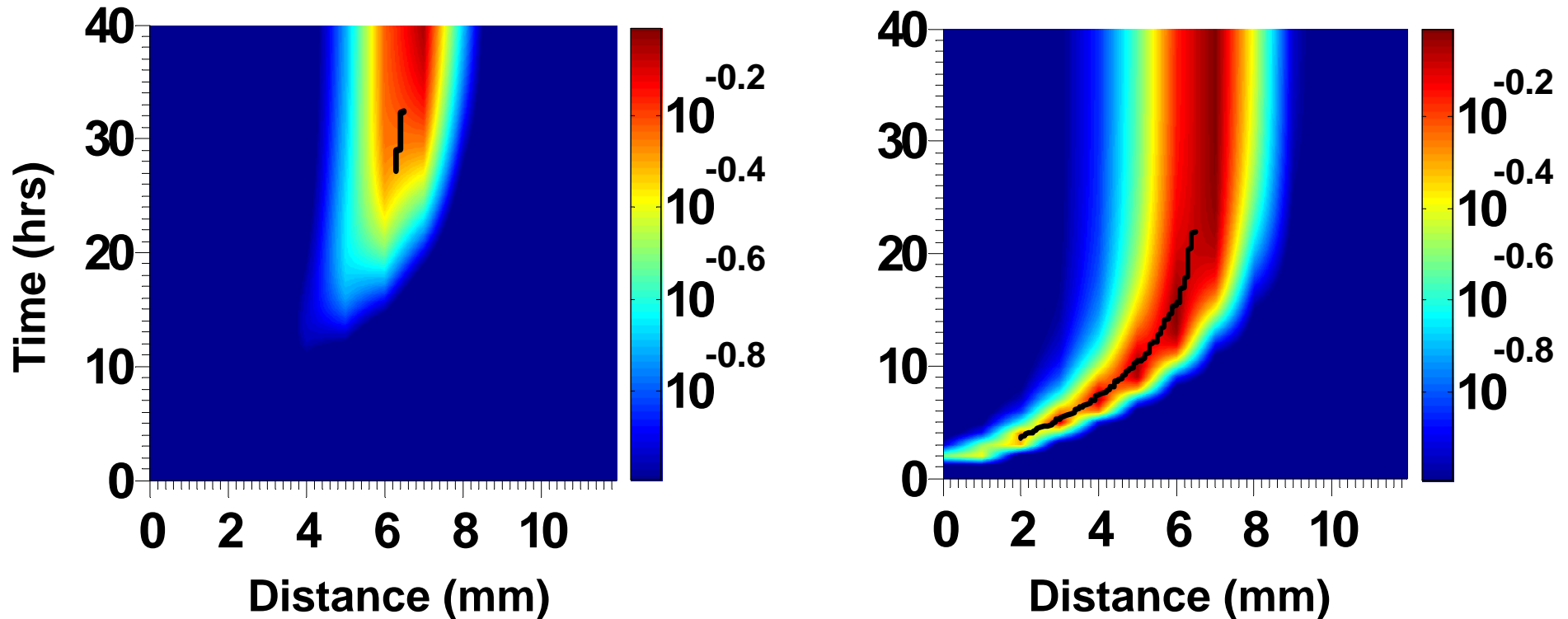
# Experimental Spatiotemporal Behavior

senders

receivers



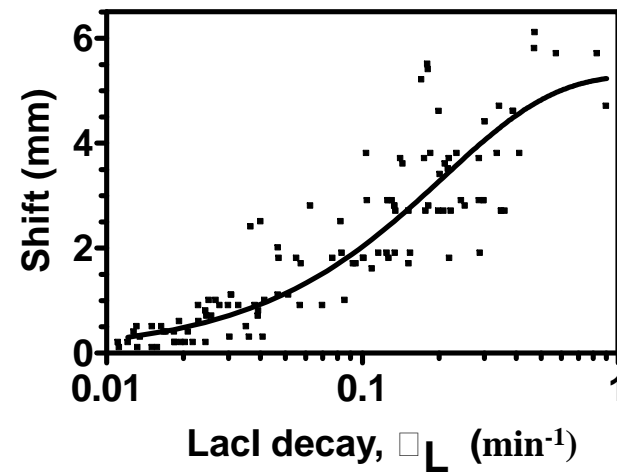
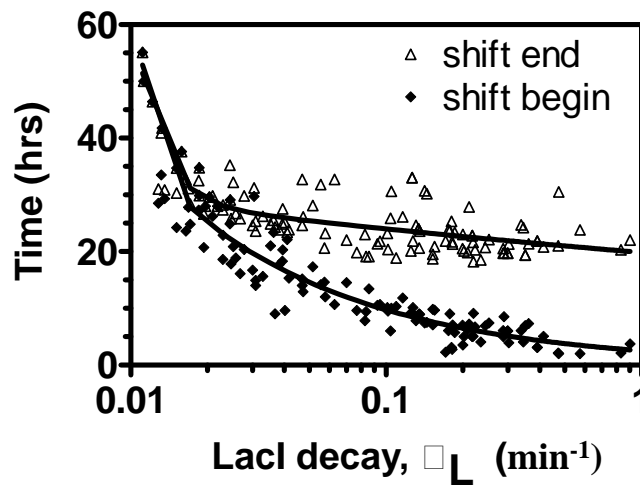
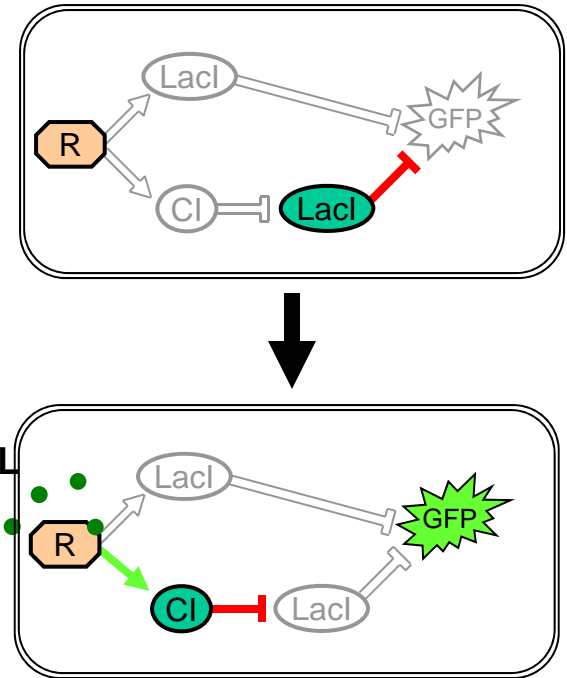
# Spatiotemporal Simulations – Shift



- Analysis of the effect of kinetic parameters on positional shift through simulations

# Regression Analysis

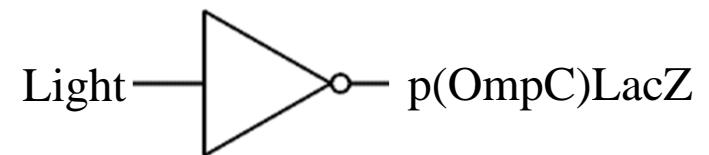
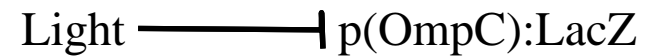
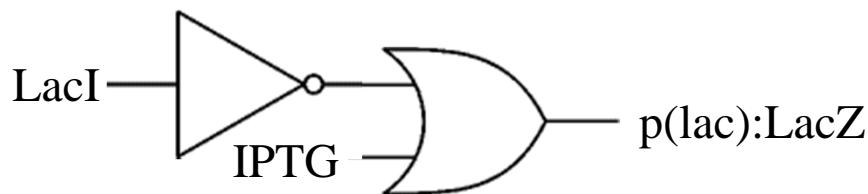
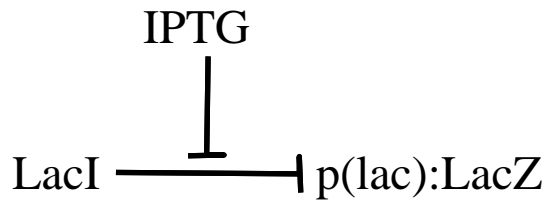
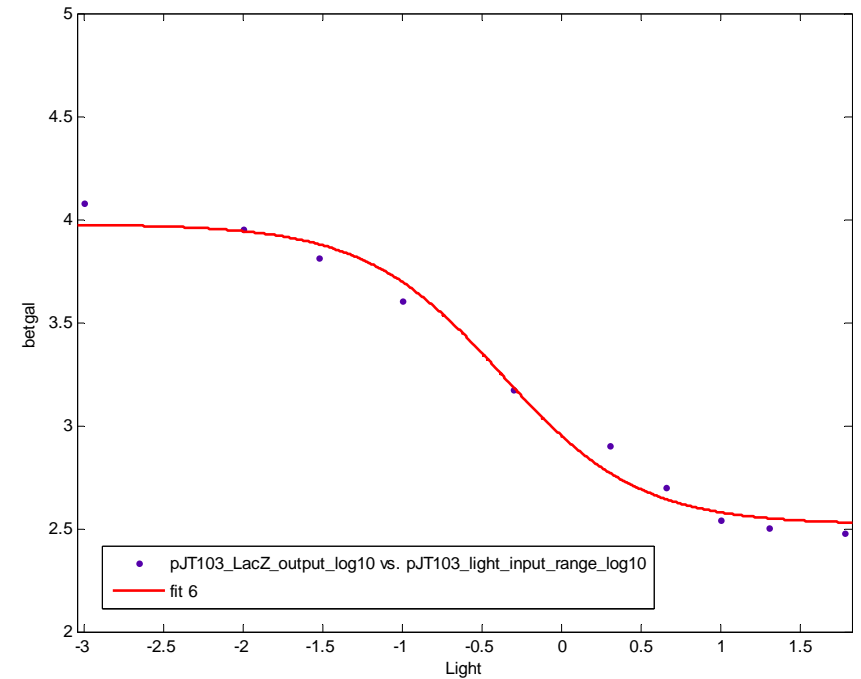
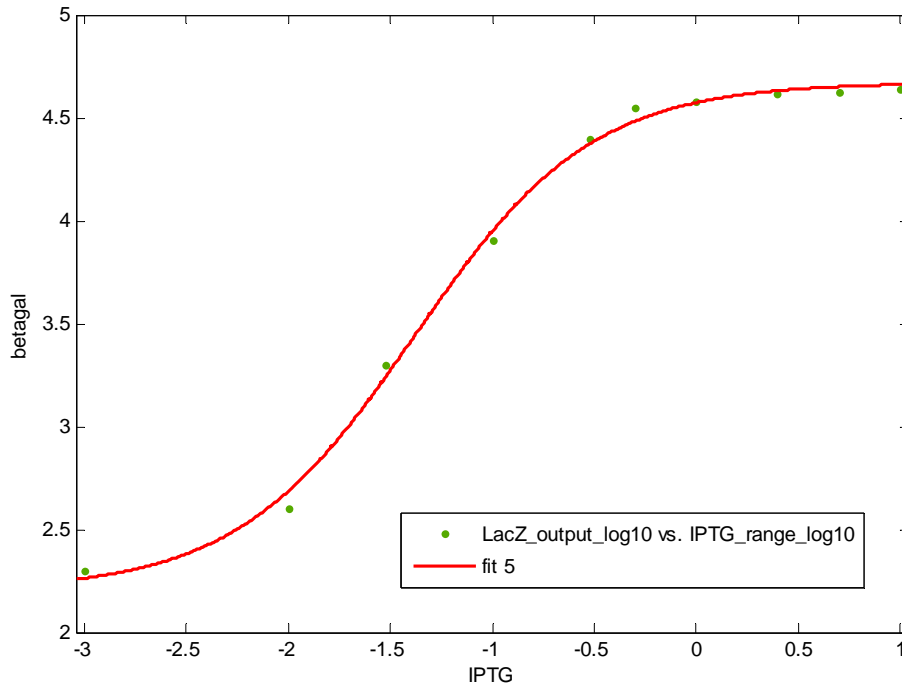
- Generated sets with random kinetic rates
- Selected for band-detect behavior (~30%)
- Regression analysis to find correlations



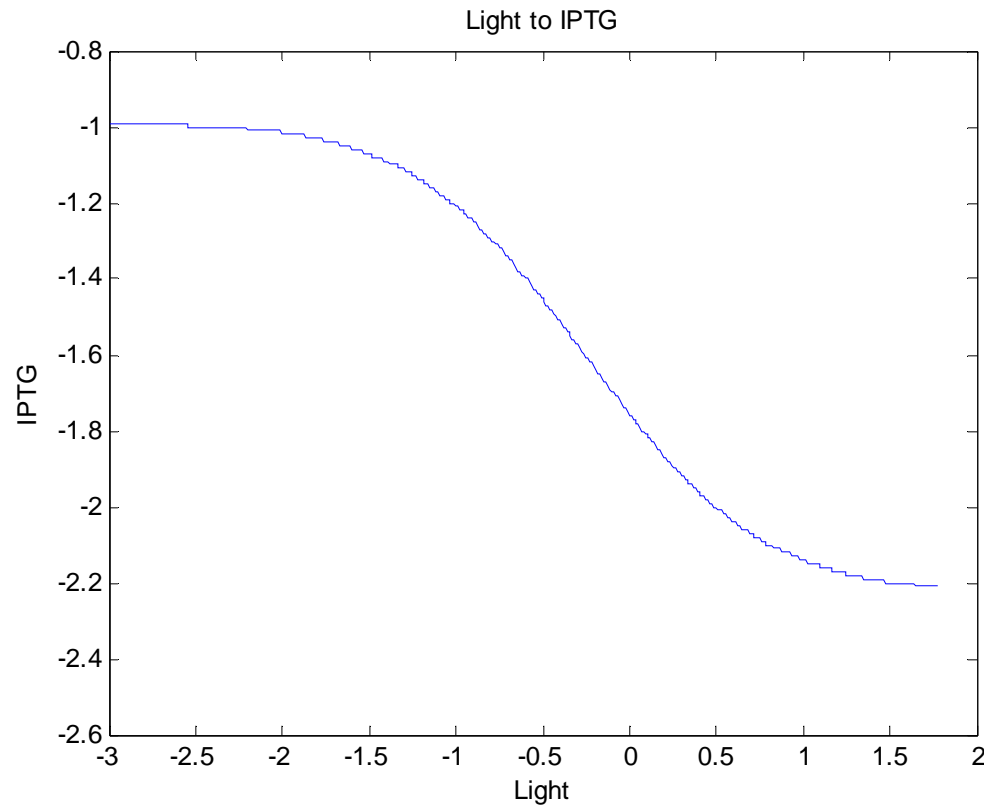


# Edge detector:

## Lab data & computational models

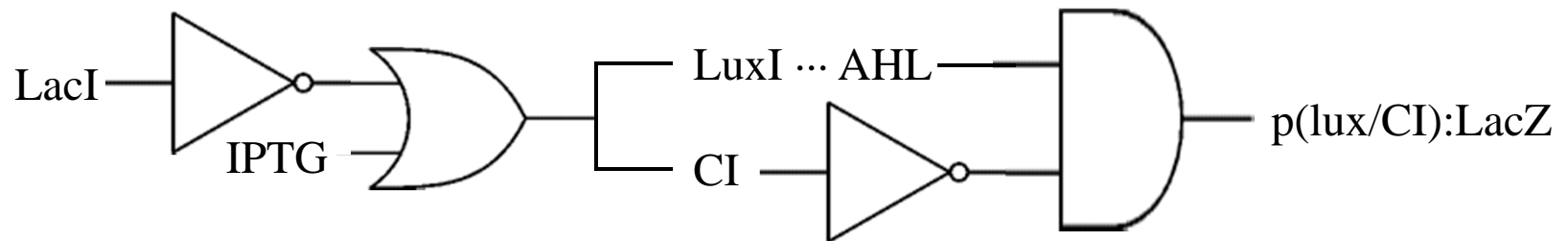
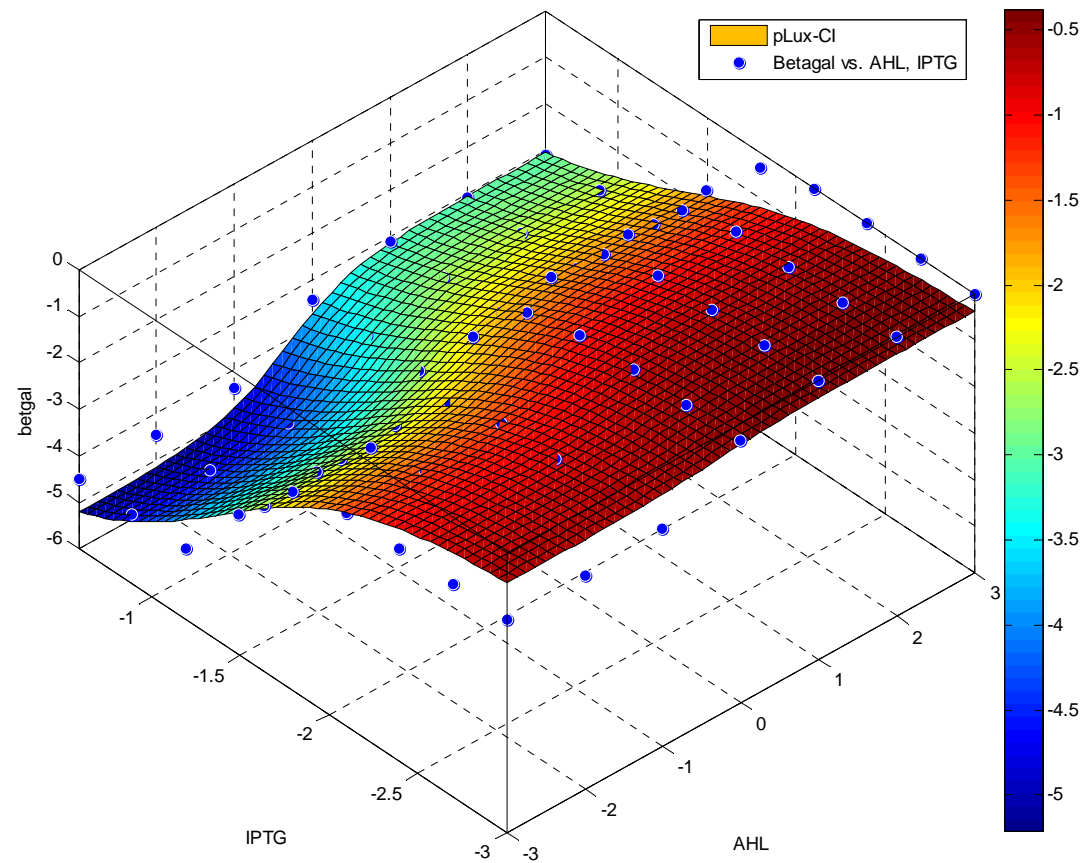


# Correlating IPTG & Light induction

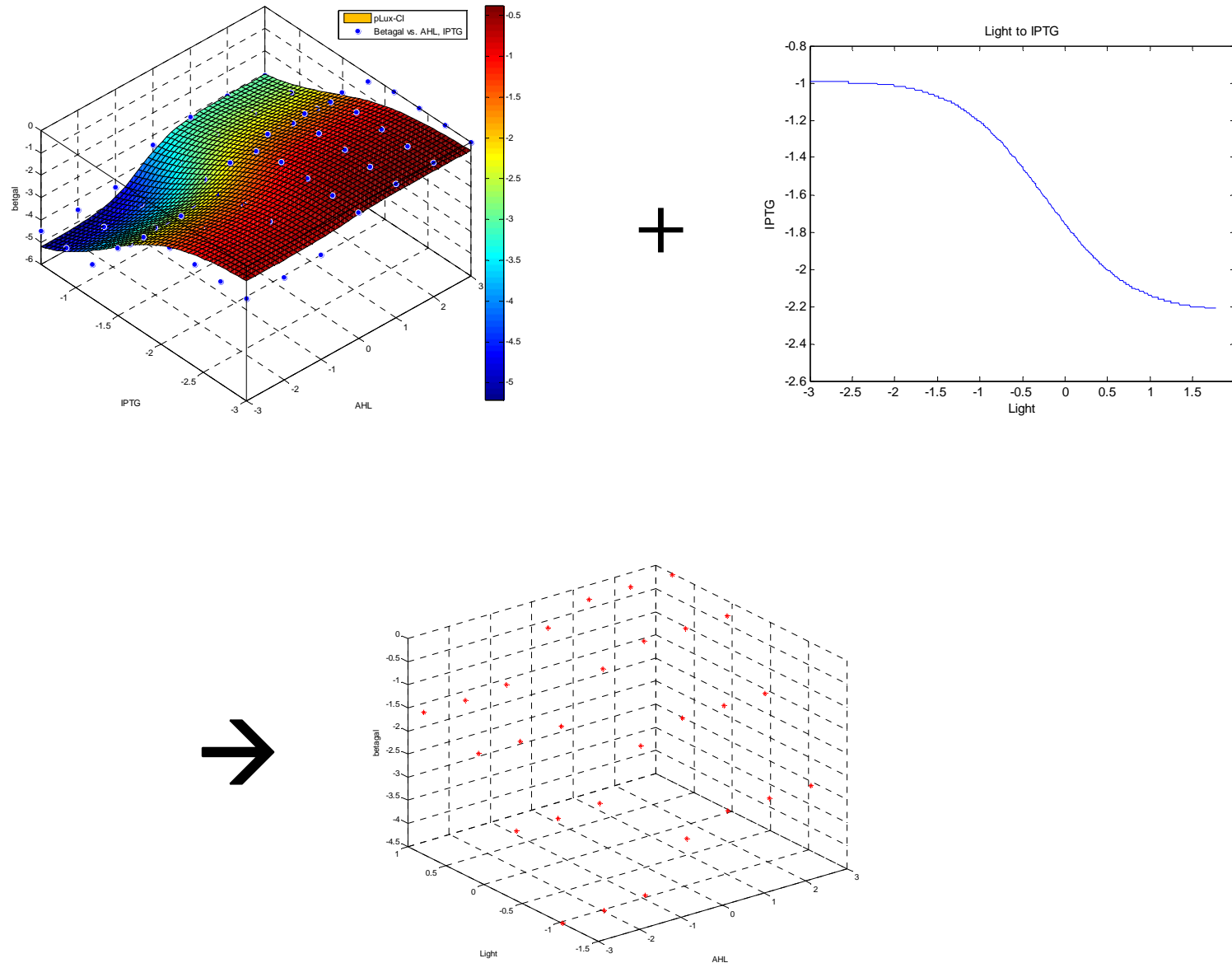


$\text{IPTG/p(lac)} \leftrightarrow \text{Light/p(Ompc)}$

# Evaluating pLux/CI with IPTG

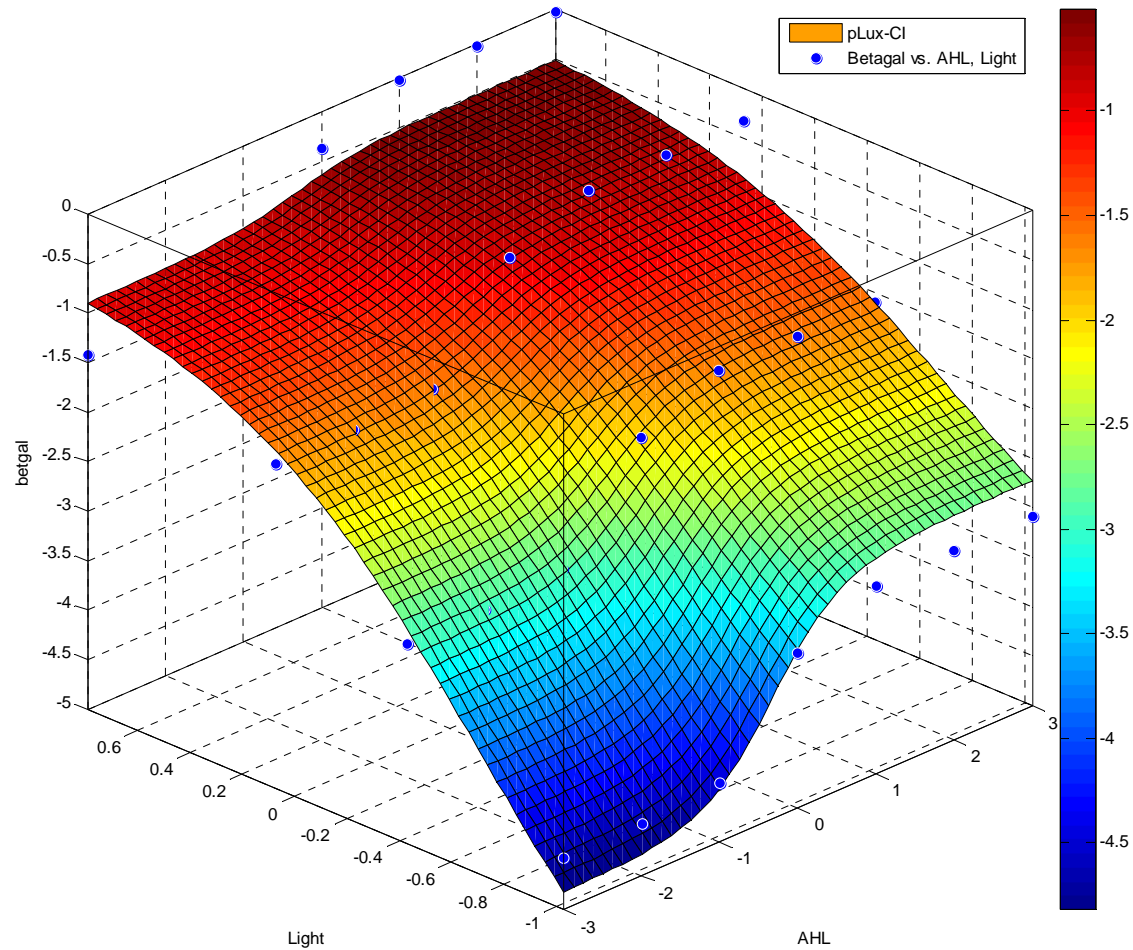


# Predicting pLux/CI with light induction





# Curve fit for pLux/CI with light induction



# pLux/CI transfer function for simulation

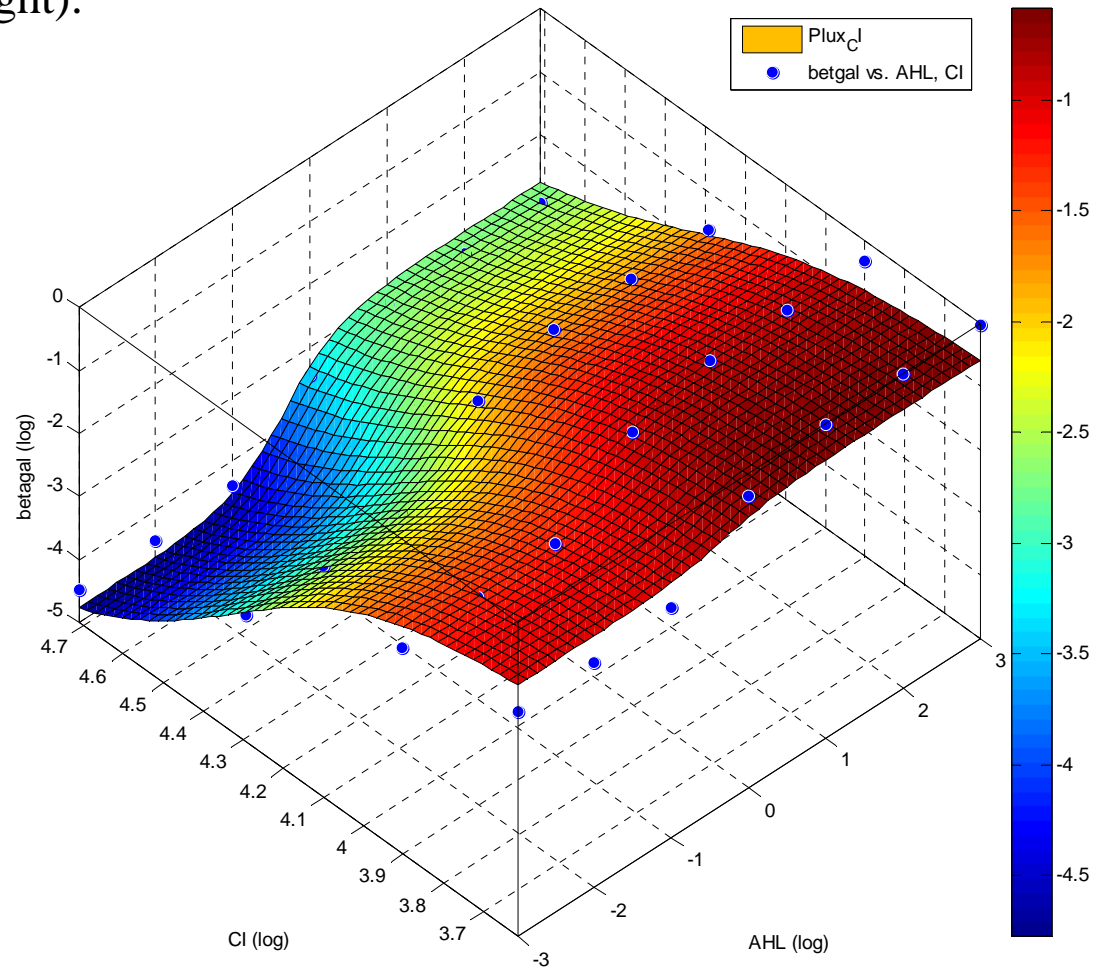
For simulation, need CI as input (not light):

$$N_{lacZ} = N_{cells} * (MU * mL * min * 0.5)$$

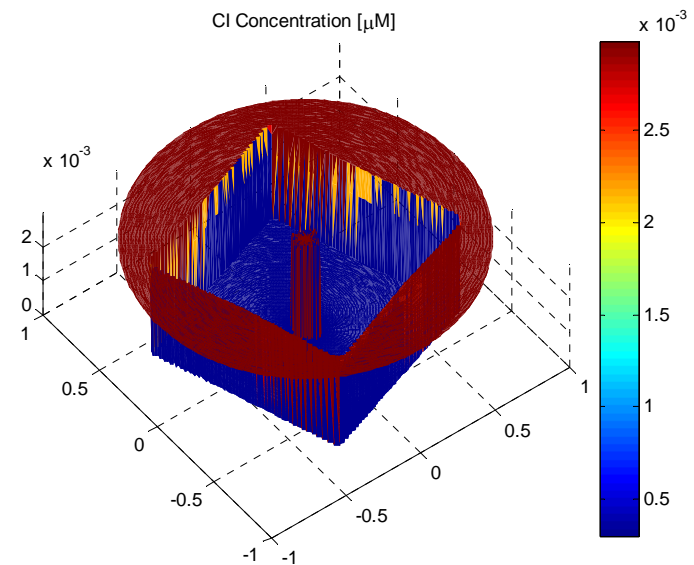
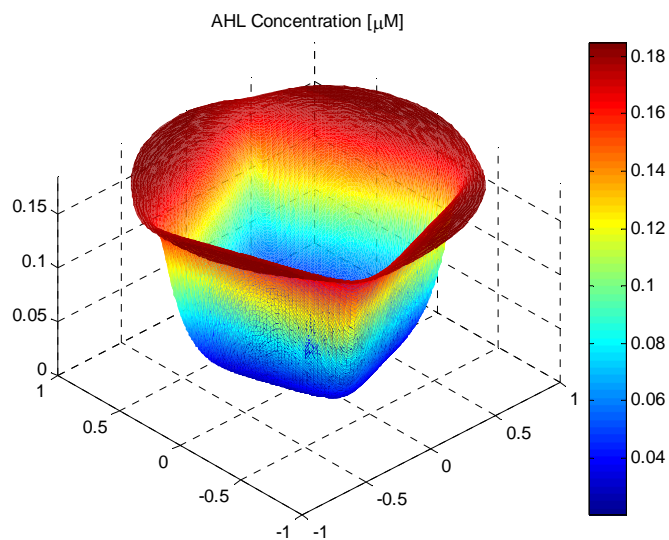
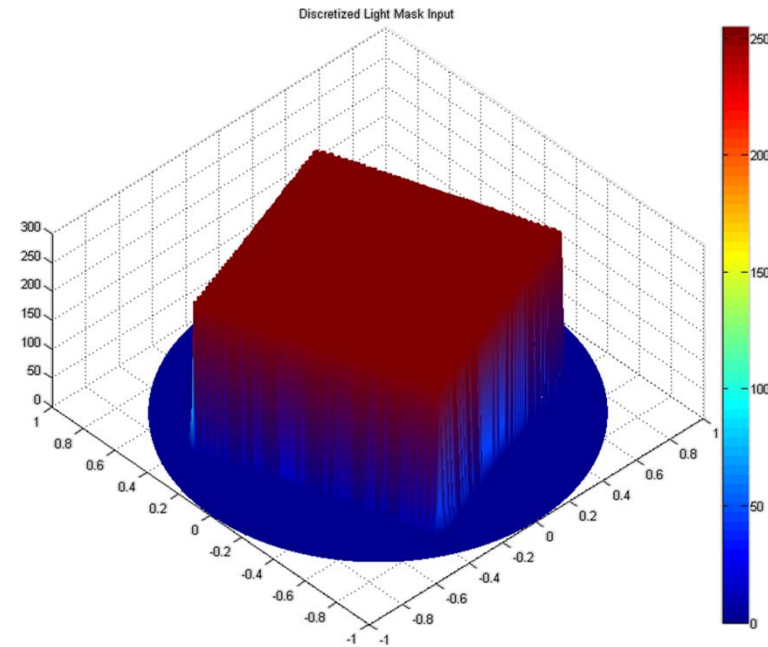
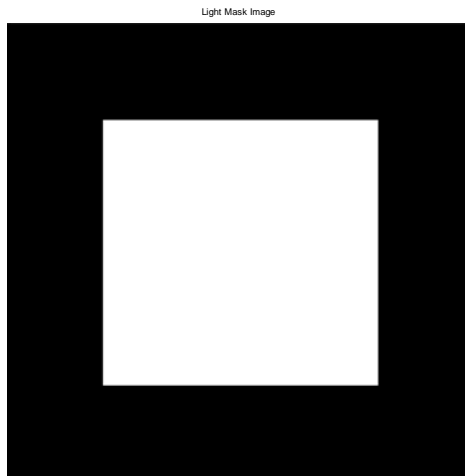
where:

- MU is the miller units
- mL is the volume of the reaction
- min is the time of the reaction

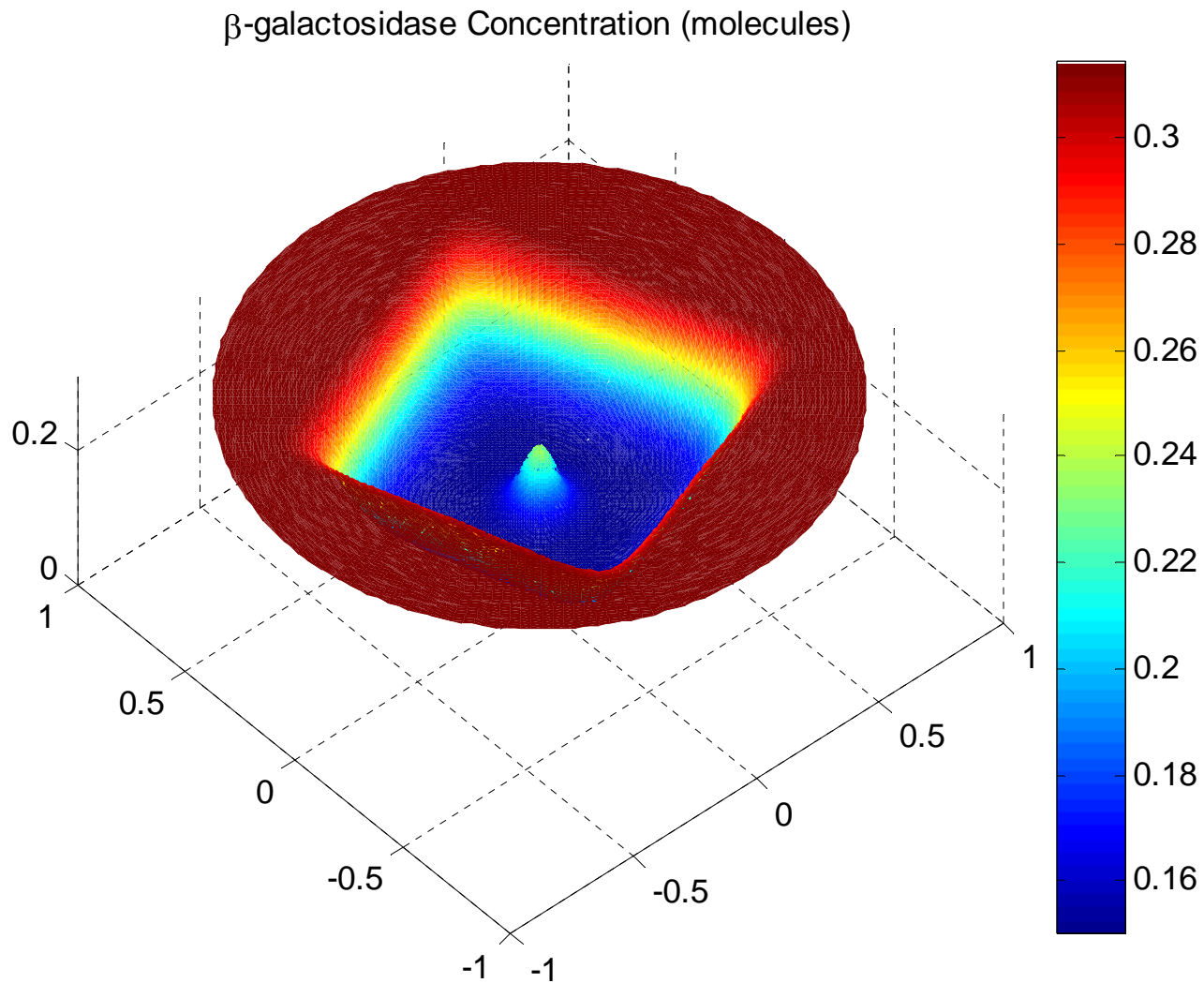
and assuming  $[CI] = [LacZ]$



# Spatiotemporal simulation: square light mask



# Simulated edge detection



Key question: with a fixed pLux/CI, will the simulations produce good edges?